

TROPICAL RAINFOREST BIRD COMMUNITY STRUCTURE IN RELATION TO ALTITUDE, TREE SPECIES COMPOSITION, AND NULL MODELS IN THE WESTERN GHATS, INDIA¹

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Studies of species distribution on elevational gradients are essential to understand principles of community organisation as well as to conserve species in montane regions. This study examined the patterns of species richness, abundance, composition, range sizes, and distribution of rainforest birds at 14 sites along an elevational gradient (500-1400 m) in the Kalakad-Mundanthurai Tiger Reserve (KMTR) of the Western Ghats, India. In contrast to theoretical possibilities, resident bird species richness did not change significantly with elevation although the species composition changed substantially (<10% similarity) between the lowest and highest elevation sites. Constancy in species richness was possibly due to relative constancy in productivity and lack of elevational trends in vegetation structure. Elevational range size of birds, expected to increase with elevation according to Rapoport's rule, was found to show a contrasting inverse U-shaped pattern because species with narrow elevational distributions, including endemics, occurred at both ends of the gradient (below 800 m and above 1200 m). Bird species composition also did not vary randomly along the gradient as assessed using a hierarchy of null models of community assembly, from completely unconstrained models to ones with species richness and range-size distribution restrictions. Instead, bird community composition was significantly correlated with elevation and tree species composition of sites, indicating the influence of deterministic factors on bird community structure. Conservation of low- and high-elevation areas and maintenance of tree species composition against habitat alteration are important for bird conservation in the southern Western Ghats rainforests.

Key words: structure and floristics, mid-domain effect, non-equilibrium dynamics, null models, community assembly

INTRODUCTION

Since the pioneering work of Terborgh (1971, 1977, 1985) and Terborgh and Weske (1975), research on altitudinal distribution patterns of birds has tended to focus on three major aspects — variation in richness with altitude, range sizes, and species turnover rates — and on one region with relatively well-documented information on bird altitudinal distributions, the Neotropics (Rahbek 1997; Patterson *et al.* 1998; Stotz 1998; Blake and Loiselle 2000). Most studies have shown a general pattern of decreasing species richness with increasing altitude, believed to mirror the well-recognised latitudinal gradient in species richness (MacArthur 1972; Patterson *et al.* 1996, 1998; Bachman *et al.* 2004). Monotonic decline in species richness with altitude may also occur when altitudinal range sizes of species increase with altitude, as suggested by Stevens (1992) in an extension of Rapoport's rule, a matter of intense debate (Rapoport 1982; Gaston *et al.* 1998; Taylor and Gaines 1999).

Other studies have suggested that species richness may show a non-monotonic, hump-shaped pattern with diversity peaking at mid-altitudes (Janzen 1973; Rahbek 1997; Stotz 1998), as in diversity-productivity gradients (Rosenzweig and Abramsky 1993). This pattern may become

apparent when data are corrected for sampling effort for local species richness (Terborgh 1977; Patterson *et al.* 1996), or area for regional species richness (Rahbek 1997; Bachman *et al.* 2004). Models that evaluate consequences of geometric constraints (hard upper and lower limits) on the altitudinal ranges of species also predict mid-altitude peaks called the 'mid-domain effect' (Rahbek 1997; Colwell and Lees 2000; Veech 2000).

In contrast to species richness, community compositional change (turnover rates) at the regional level may increase with altitude (Rahbek 1997), or show peaks and troughs corresponding to transition zones between lowland and montane avifauna (Patterson *et al.* 1996; Stotz 1998). Little is known of factors influencing bird community composition at local levels in tropical rainforests. Along successional gradients, habitat structure and tree species composition influence bird community structure, wherein structurally and floristically similar sites have more similar bird communities (Raman *et al.* 1998). Along altitudinal gradients, past studies of local bird community structure have largely ignored the relative influence of deterministic differences between sites in habitat attributes versus the effects of chance, null models, or altitude *per se*.

The present study explores variation in local bird

community structure, species composition, and turnover rates along an altitudinal gradient in a tropical rainforest of the Western Ghats of India. Although recognised as a global biodiversity hotspot (Myers *et al.* 2000) and an endemic bird area (Stattersfield *et al.* 1998), this region is peculiar in being relatively depauperate in its rainforest bird community due to historical isolation from larger tracts of rainforest in northeast India and southeast Asia (Daniels *et al.* 1992). Here, we explore how bird community attributes vary with altitude in tropical rainforest. Specifically, we ask: does species richness show a monotonic decline with altitude or a hump-shaped pattern due to the 'mid-domain effect'? Is there support for Rapoport's rule? Does altitudinal distance and degree of change in habitat influence and correlate with change in bird community composition or are turnover rates estimated accurately by null models that simulate non-equilibrium dynamics? The results are used to assess the relative influence of deterministic *versus* non-equilibrium factors on tropical rainforest bird community structure.

MATERIAL AND METHODS

Study area

The 1600 km long chain of hills called the Western Ghats runs along the west coast of the Indian peninsula from 8° N to 21° N. Moist forests, including tropical wet evergreen forest, are found largely south of 16° N (Pascal 1988), and contain a higher diversity of endemic plant and animal taxa (Ali and Ripley 1983; Daniels 1992; Kumar *et al.* 2004). The fieldwork was carried out in the Kalakad-Mundanthurai Tiger Reserve (KMTR, 895 sq. km, 8° 25' to 8° 53' N, 77° 10' to 77° 35' E) in the Agasthyamalai region at the southern end of the Western Ghats. KMTR ranges between 50 m and 1700 m above msl with rainforest occurring chiefly above 500 m. This reserve along with adjoining areas has one of the largest remaining contiguous tracts (over 400 sq. km) of relatively undisturbed tropical rainforest in the Western Ghats (Ramesh *et al.* 1997). The rainforest vegetation in KMTR has been classified mid-elevation tropical wet evergreen rainforest of the *Cullenia exarillata-Mesua ferrea-Palaquium ellipticum* type (Pascal 1988; Ganesh *et al.* 1996; Parthasarathy 2001). Within the rainforest, the mean monthly daytime temperature ranges between 19 °C in January and 24 °C in April-May (at Sengaltheri, 1040 m, range 15-31 °C). The total rainfall was 2283 mm in 1998 and 2230 mm in 1999 at Sengaltheri. There are three seasons: (a) dry season (February to May), (b) southwest or summer monsoon (June to September), and (c) northeast or winter monsoon (October to January). KMTR receives over half its annual precipitation during the northeast monsoon.

Although 278 bird species have been recorded in and around KMTR, only 84 species occur in rainforests including 12 endemics and 13 winter migrants (Raman 2001). A majority of species breed between late January and May (Ali and Ripley 1983, TRSR pers. obs). The sites chosen for intensive bird community sampling were located at Kannikatti (740 m, 8° 37' N and 77° 16' E), Sengaltheri (1040 m, 8° 31' N and 77° 26' E), and Kakachi (1220 m, 8° 33' N and 77° 24' E) in KMTR. These areas contain rainforests between 500 m and 1400 m altitude, contiguous with rainforests elsewhere in the Reserve. Fourteen sites located along this altitudinal gradient in relatively undisturbed rainforest were selected randomly, within logistical constraints, around three forest camps (four accessed from Kannikatti, six from Sengaltheri, and four from Kakachi) and located on topographic maps using landmarks and a global positioning system receiver (GPS).

Bird surveys

The fixed-radius point count method was used to survey bird populations in each site in a relatively uniform and efficient manner during the main breeding season (Verner 1985; Ralph *et al.* 1995). Point count surveys (5 min duration) were carried out during the first three hours after sunrise (see Raman 2003 for details of field technique). Densities estimated by the fixed radius approach were used as they were highly correlated to variable-radius point count estimates across species (Raman 2003).

Each sampled site represented an area of around 12-15 ha lying at the designated altitude and was surrounded by contiguous rainforests on at least three sides. Within each site a 600-700 m long transect or narrow animal trail was marked at 25 m intervals for the point count surveys. At each site, 25 point count surveys (yielding 167-247 detections and an estimated 334-597 individual birds per site) were carried out, excluding the Kodayar site where only 18 counts were carried out. Sampling of all sites around Sengaltheri and Kakachi was carried out mostly between February and May 1998, during the peak breeding season when resident and migrant birds were present. For reasons of logistics, the four sites around Kannikatti were sampled only in March 1999.

Vegetation sampling

In each of the 14 sites, densities of trees greater than 30 cm girth at breast height (GBH at 1.3 m) were estimated using the point-centred quarter method (PCQ, Krebs 1989). Thirteen PCQ plots, with successive plots spaced 50 m apart, were measured in each site, giving a sample of 52 trees per site. Comparison with two completely enumerated 1 ha plots (>1800 stems each) in Sengaltheri showed that the density estimates from this PCQ sample was <2% different from the

more laborious enumeration. All trees were identified to species, or in a few cases to genus (Gamble and Fischer 1915-1935; Pascal and Ramesh 1997). Distance from plot centre to the middle of the bole and GBH were recorded for each tree. At each of the 13 PCQ plots, 2 m radius plots were laid to enumerate shrubs. In addition, the number of cane plants and the presence or absence of bamboo within 5 m radius was recorded.

Altitude, canopy measures and leaf litter variables were measured at 25 points, evenly spaced 25 m apart, in each site. Canopy height was measured using a rangefinder and percentage canopy cover using a spherical densiometer. Vertical stratification was assessed at these 25 points by noting the presence or absence of foliage in the following height intervals (in metres): 0-1, 1-2, 2-4, 4-8, 8-16, 16-24, 24-32, and >32, directly above and in a 0.5 m radius around each point (Raman *et al.* 1998). Leaf litter depth on the forest floor was measured using a calibrated wooden probe. As ground vegetation and litter were disturbed along trails, the samples were taken 10 m away from trails into the rainforest interior.

Data analysis

The cumulative list of bird species recorded in each site was used as a basic measure of bird species richness. However, because only 18 point counts were sampled at the Kannikatti site, we obtained standardised estimates of bird community parameters for all sites for 17 sampled points. Using the program EstimateS (Colwell 1997), 100 permutations (17 samples without replacement) were performed to estimate the following parameters and their standard deviations: (1) bird species richness, measured as the cumulative number of species; (2) bird abundance, estimated as the number of individual birds/ha; (3) Shannon-Weiner diversity, calculated as $-\sum(p_i \ln p_i)$, where p_i is the proportion of the i^{th} species and the summation is across all species in the pooled samples; (4) rarefaction richness, standardising for sampling effort using Coleman rarefaction curves (Colwell 1997); (5) bootstrap species richness, an incidence-based estimator denoted as $S_{boot} = S_{obs} + \sum(1 - p_k)^m$, where p_k is the proportion of m samples with species k and the summation is across all S_{obs} species in the pooled samples; and (6) Chao 1 richness, an abundance-based estimator of species richness (see Colwell 1997 for the formula and correct variance estimator of this index).

Smaller scale point richness and abundance estimates were also obtained by averaging across replicate point count surveys in each site. The average number of species, detections, and individuals per point and their standard errors were estimated for all species combined and for resident species alone. For each site, tree density and basal area were calculated using the PCQ method (Krebs 1989). Average values across replicate sampling points in each site were calculated

for other vegetation and site variables: shrub, cane, and cardamom densities, leaf litter depth, canopy height and cover, and altitude. Vertical stratification was measured as the average number of strata with foliage across the 25 points sampled in each site. The coefficient of variation of vertical stratification was used as an index of horizontal heterogeneity (following Raman *et al.* 1998). The total number of tree species recorded in the PCQ plots was recorded as a measure of tree species richness. Relationships between bird community and vegetation variables were assessed using Kendall rank correlations (Siegel and Castellan 1988).

Altitudinal range size of each species was estimated as the difference between the maximum and minimum altitude at which the species was recorded in the point count surveys. The altitudinal range midpoint and a weighted range midpoint, estimated using the abundance of each species in each site, were calculated for each species. Similarities in bird community, foliage profile, and tree species composition between sites were computed using the Morisita index that is least sensitive to sample size effects (Wolda 1981; Raman *et al.* 1998). The matrix of pair-wise dissimilarities (100-Morisita index in %) in bird community composition was related to corresponding matrices of similarity/distance in tree species composition, foliage profile, altitude, and geographic distance between sites using Mantel tests (Manly 1994). Geographic distance between sites was measured as the straight-line distance between sampling sites. Statistical significance of Mantel tests was assessed through 10000 random permutations. To examine the independent effects of different variables, partial Mantel tests, derived from the Kendall tau approach, were used (Hemelrijk 1990). The tree and bird community similarity matrices were also used to ordinate study sites using multidimensional scaling (Manly 1994).

We used correlation and regression techniques to assess statistical significance of altitudinal trends. Linear and non-linear (chiefly quadratic) regressions were applied to describe spatial trend data. We also used non-parametric Kendall (*tau*) rank-order correlation coefficients to assess statistical significance (Siegel and Castellan 1988).

Null model analyses

The mid-domain effect (Colwell and Lees 2000) was assessed using the null model of Veech (2000). Species' altitudinal range widths were retained but located randomly on the altitudinal gradient between 500 and 1400 m (divided into 18 zones of 50 m width) in 1000 simulations to obtain an expected species richness curve along this gradient. The observed pattern of species richness was compared to the simulated curve to obtain a mean displacement, D , where $D = (\sum d_j)/18$, where d_j is the absolute difference in species

richness between the two curves being compared for each of the 18 zones (Veech 2000). A further 1000 simulations of the data were used to determine the distribution of D values in each run and to assess statistical significance.

Null models were also used to assess the likelihood of obtaining, under the null model assumptions, estimates equivalent to two measures of the average similarity between sites in bird community composition. These measures were: (i) average similarity between all possible pairs of sites ($N = 91$ comparisons), and (ii) average similarity between successive sites along the altitudinal gradient ($N = 13$ comparisons). The Jaccard index (Krebs 1989) was used to measure similarity in bird community composition. With the presence-absence matrix of species across sites, five null models were evaluated:

1. Unconstrained model: matrix total constant, row and column totals varied;
2. Saturation model: fixed column totals (species richness at each site);
3. Distribution model: fixed row totals (number of sites each species occurred in);
4. Doubly constrained model: fixed row and column totals by a procedure equivalent to the Knight's tour algorithm (Sanderson 2000);
5. Range-contiguity model: this model simulated, in addition to the distribution constraint, the contiguity of ranges of species along the gradient. Simulations showed that species occurrences were distributed more contiguously along the gradient than expected by random placement ($P < 0.001$). Using the observed occurrences in the species-by-site matrix, the relative probability of occurrence in any one of the 12 intermediate sites (excluding the 2 'edge' sites—the lowest and highest site) was found to be enhanced by a factor of 9.0 if the species also occurred in sites immediately above and below, and by a factor of 4.5 if it occurred only in one adjacent site either above or below. These enhancement factors were in relation to the probability of occurrence in a site when the species was absent from both adjacent sites on the gradient. For the edge sites, occurrence was enhanced by a factor of 3.6 if the species occurred in the adjacent site in relation to occurrence when it was absent in the adjacent site. These were then used to distribute occurrences in simulations to produce probabilistic contiguous distributions.

These models attempt to simulate the effects of chance and biological constraints on the assembly of species communities (Farnsworth and Ellison 1996; Stone *et al.* 1996; Sanderson 2000). In contrast to the unconstrained model, the saturation, distribution, and doubly-constrained models implement constraints on the number of species that can fill each site and simulate the effects of interspecific differences

in distribution and occurrence probabilities. Our range-contiguity model incorporates, in addition, constraints on species' altitudinal distributions and contiguity of ranges. We retained in the contiguity model the number of intermediate absences or 'holes' in range but randomised their location. Even with these constraints, we checked and established that over 5000 different matrices existed in simulations. For each of the above models, we simulated 5000 randomised matrices and the averages of the obtained distribution of similarity values (between all possible pairs of sites and adjacent sites) were statistically compared to the observed average similarity values.

RESULTS

Bird species richness, diversity, and abundance

Across the 14 sites, we obtained over 2900 detections comprising approximately 6600 individuals of birds belonging to 67 species. Of these, nine species were latitudinal (winter) migrants and the remaining were resident species. Thirteen species were detected only once and a further seven species were seen less than five times overall. Total bird species richness (cumulative list in each site) was significantly negatively correlated to altitude of the site (Kendall rank-order correlation, $\tau = -0.52$, $n = 14$, $P = 0.01$, Table 1). Resident bird species richness appeared to decline with altitude, but this trend was not significant statistically ($P = 0.08$). The standardised estimates for 17 point counts obtained over 100 runs (sampling with replacement) of species richness, Shannon-Weiner diversity, Coleman rarefaction richness, bootstrap richness, and the Chao1 richness were all significantly negatively correlated to altitude ($P < 0.05$, Table 1). Most of the decrease occurred between 500 m and 900 m altitude, after which there was little change in species richness. Bird abundance showed a converse positive association with altitude that was marginally significant ($\tau = 0.39$, $n = 14$, $P = 0.055$, Table 1). The per point estimates of total and resident bird species richness, detections, and abundance showed significant variation across the 14 sites (one-way ANOVA, $F_{13,329} > 5.58$, $P < 0.004$). Whereas the number of detections per point showed weak negative correlations with altitude ($\tau = -0.41$, $P = 0.04$), the trends for total and resident species richness and detections with altitude were non-significant. Bird abundance per point increased with altitude, weakly and non-significantly in the case of total abundance ($\tau = 0.38$, $P = 0.06$) and significantly in the case of abundance of resident species ($\tau = 0.52$, $P = 0.009$).

Vegetation and birds: correlations

Vegetation variables were not significantly correlated

Table 1: Bird community parameters along an altitudinal gradient in the rainforests of Kalakad-Mundanthurai Tiger Reserve, India, showing Kendall rank-order correlations with altitude

Transect Code	A	B	C	D	E	F	G	H	I	J	K	L	M	N	Correlation with altitude		
Mean altitude (m)	558	646	751	843	900	980	1092	1228	1256	1259	1265	1287	1341	1359	t	Z	P
Cumulative species richness																	
Total species	40	36	36	34	30	29	31	29	32	31	30	25	28	32	-0.520	-2.59	0.0096
Resident species	35	30	30	29	26	25	27	26	30	27	28	25	25	30	-0.352	-1.75	0.0798
Standardised estimates (Estimates)																	
Bird species richness	36.8	33.0	33.0	31.1	27.4	25.9	28.2	27.0	28.9	27.5	26.3	21.9	27.6	29.97	-0.407	-2.03	0.0428
SD	1.60	1.46	1.60	1.34	1.55	1.99	1.38	1.31	1.37	1.83	1.90	1.72	0.60	1.15			
Bird abundance (individuals/ha)	22.6	24.3	21.7	22.3	23.0	17.0	22.1	29.7	24.8	25.5	20.7	25.2	25.8	30.5	0.385	1.92	0.0554
SD	0.81	1.10	1.20	1.18	1.64	1.51	1.61	1.50	1.63	1.58	1.48	2.15	0.68	1.93			
Shannon-Weiner diversity	3.22	3.03	3.06	3.02	2.73	2.74	2.82	2.69	2.79	2.63	2.48	2.07	2.53	2.8	-0.626	-3.12	0.0018
SD	0.04	0.04	0.05	0.04	0.08	0.08	0.05	0.07	0.06	0.07	0.11	0.09	0.03	0.05			
Coleman rarefaction richness	37.7	34.1	34.2	32.3	28.5	27.8	29.8	28.1	31.4	29.1	27.9	23.4	27.8	31.3	-0.538	-2.68	0.0073
SD	1.29	1.18	1.16	1.14	1.06	0.99	0.96	0.84	0.72	1.23	1.25	1.10	0.41	0.76			
Bootstrap richness	41.1	37.1	37.3	34.3	30.8	29.6	31.5	29.8	32.3	31.2	30.4	25.2	30.5	33.1	-0.429	-2.14	0.0328
SD	2.11	1.92	2.15	1.79	1.98	2.45	1.80	1.71	1.86	2.42	2.46	2.19	0.76	1.42			
Chao1 richness	40.6	33.9	35.1	32.1	28.3	26.2	28.5	27.2	29.0	28.3	28.3	23.6	28.3	30.2	-0.420	-2.09	0.0365
SD	6.31	2.06	4.38	2.08	2.20	0.86	1.19	0.19	0.46	1.60	3.70	4.38	2.08	1.00			

with altitude (Kendall correlations, $P > 0.05$), except for bamboo culm density ($\tau = 0.52, n = 14, P = 0.01$) mainly due to the occurrence of bamboo at only two higher altitude sites. The six standardised estimates of bird species richness, Shannon-Weiner bird diversity, and the per-point total and resident bird species richness were all significantly negatively correlated to one of the vegetation variables: leaf-litter depth ($\tau < -0.45, n = 14, P < 0.03$ in all cases). These variables were also not correlated to any of the other vegetation variables. Bird abundance (standardised and per-point) and the total number of bird detections per point were not correlated to any vegetation variable. The average number of resident bird

detections per point was negatively correlated to leaf litter depth ($\tau = -0.47, P = 0.037$).

Range distributions, sizes, and species richness

The range size and distribution pattern across the 58 resident bird species varied from species occurring in one to all 14 sites spanning the entire gradient. Twenty-three species (39.7%) were either restricted to lower altitudes or showed declining trends with altitude in population density (Kendall correlations, $P < 0.10$, Appendix). Sixteen species (27.6%) displayed a trend of increase in population density with altitude ($P < 0.10$) or were restricted largely to higher altitude sites (Appendix). Only two species of woodpeckers, *Dinopium javanense* and *Dryocopus javensis*, appeared to be restricted to mid-altitudes. The spatial distributions of the remaining species were widespread and relatively uniform (14 species) or could not be determined due to low sample sizes (3 species).

Across the gradient, the number of resident bird species whose ranges intercepted each 50 m altitude zone was nearly constant at around 35 species (Fig. 1a). This was significantly different from the Veech (2000) null model that predicted a mid-domain peak (Fig. 1a, $d = 6.51, P < 0.001$). The relationship between altitudinal range size of a species and the weighted midpoint of its altitudinal range was also non-linear and quadratic (Fig. 1b, fitted curve: $y = -0.0046x^2 + 8.8633x - 3496.8, R^2 = 0.72$). Similarly, in contrast to the expectation under Rapoport's rule, the mean altitudinal range of species in each site showed a quadratic relationship with the altitude of the site (Fig. 1c, fitted curve: $y = -0.0005x^2 + 8.8839x + 216.99, R^2 = 0.78$).

Bird species turnover

Species turnover was considerable along the altitudinal gradient. Non-metric multidimensional scaling ordination using the Morisita dissimilarity matrix of the data on tree and bird species composition showed similar patterns (stress = 0.070 and 0.034, Fig. 2a, b, respectively). Sites below 900 m tended to cluster together as did sites above 1100 m, while intermediate sites were relatively dissimilar to the others (with exceptions shown in tree species composition by site D at 843 m and site M at 1341 m).

Partial Mantel tests showed that dissimilarity in bird community composition between sites was independently positively correlated to both altitudinal distance between sites ($T = 0.60, P < 0.0001$, Fig. 3a) and dissimilarity in tree species composition ($T = 0.22, P = 0.013$, Fig. 3b). Correlations with both these variables were significant even when controlled for the effects of geographic distance ($T = 0.69$ and $0.51, P < 0.0003$). In contrast, geographic distance between sites had no effect on bird species composition, when the effects of

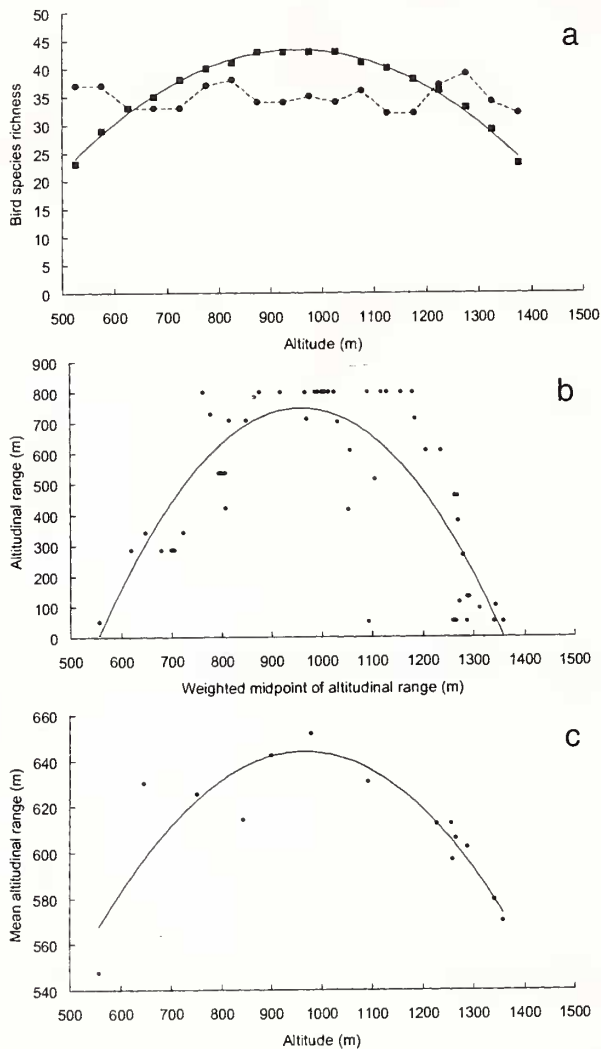


Fig. 1: Bird species richness and range size relationships in rainforests of KMTR; (a) upper panel: evaluation of the Veech (2000) null model of the mid-domain effect—line shows best fit curve to null model estimates against observed curve (dotted line); (b) middle panel: altitudinal range size of 58 resident bird species in relation to the weighted mid-point of their altitudinal range; and (c) lower panel: mean altitudinal range of species at each of the 14 sampled sites in relation to altitude of the sites.

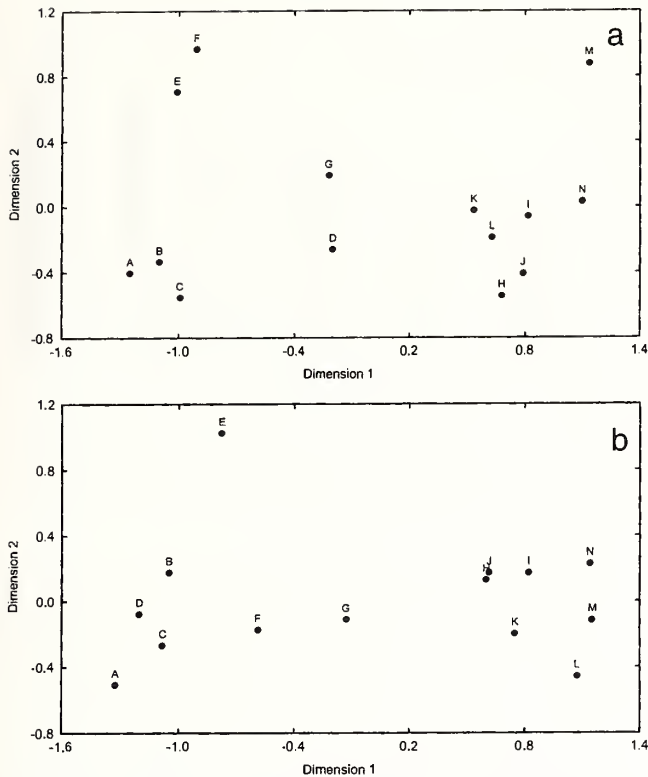


Fig. 2: Ordination of study sites on the basis of similarity in (a) tree species composition (upper panel) and (b) bird community composition (lower panel) using multidimensional scaling. Sites are named A to N in increasing order of altitude.

altitudinal distance were controlled for ($T = 0.08$, $P = 0.12$), indicating the primary importance of altitude. We found no significant effect of foliage profile dissimilarity on bird community composition.

Species turnover between sites (average of 91 all-possible pairs) was 0.529 ± 0.139 SD by the Jaccard similarity index and 0.608 ± 0.251 SD by the Morisita similarity index. This was lower than average similarity in bird community composition between adjacent sites (Jaccard index: 0.693 ± 0.107 SD, Morisita index: 0.860 ± 0.103 SD). The pattern of turnover was not monotonic or increasing with altitude, and remained relatively low but for higher turnover (lower similarity) around 1000 m (Fig. 4).

Null models of species turnover

We assessed whether the five null models of bird species turnover could accurately estimate the observed Jaccard similarity between all possible pairs of sites (0.529 ± 0.139 SD) and between adjacent pairs of sites (0.693 ± 0.107 SD). The unconstrained model and saturation models gave similar results, producing similarity estimates that were significantly lower than the observed ($P < 0.0002$, Fig. 5). The estimated average similarities between all possible pairs of sites by the range-contiguity model and the double-constraint

model were close to that actually observed (0.516 vs. 0.529), but were, nevertheless, significantly lower ($P < 0.0002$). The double-constraint model also performed poorly in estimating average similarity between adjacent sites (0.517 vs. 0.693). The range-contiguity model estimated similarities between adjacent sites that were higher and much closer to the observed value (0.622 vs. 0.693). However, the model estimates were significantly lower than the observed values for double-constraint and range contiguity models ($P < 0.0002$, Fig. 5).

DISCUSSION

Species richness

Species richness of resident rainforest birds varied little despite substantial change in community composition across the altitudinal gradient, in contrast to expectation from the four main models hitherto proposed (Rahbek 1997). The general model, propounding a monotonous decline in species richness with altitude in parallel with an assumed decline in productivity, finds weak support, if any, when all bird species including latitudinal migrants are considered. The pattern of range sizes observed in this study results in local species richness patterns that depart significantly from expectations

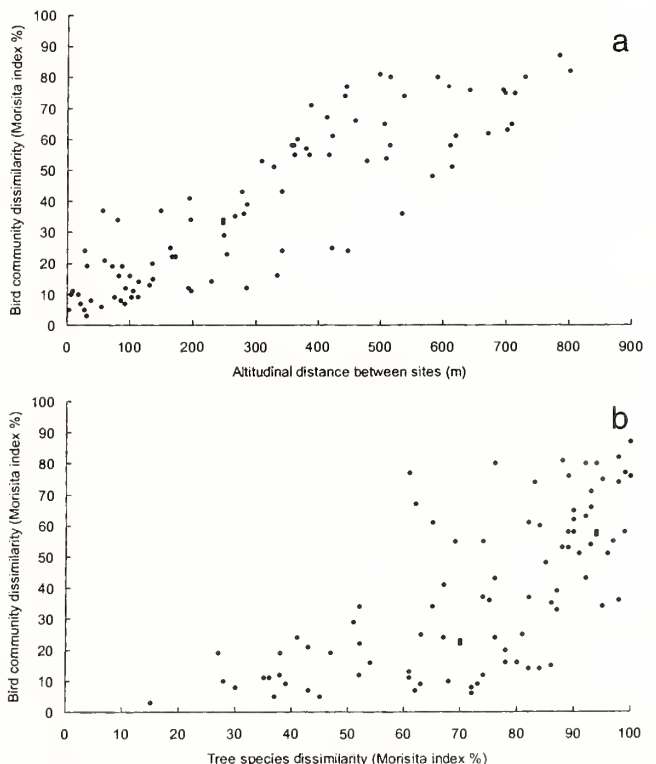


Fig. 3: Bird community dissimilarity between sites (91 all-possible pairs) in relation to corresponding between-site (a) altitudinal distance (upper panel) and (b) tree species dissimilarity (lower panel).

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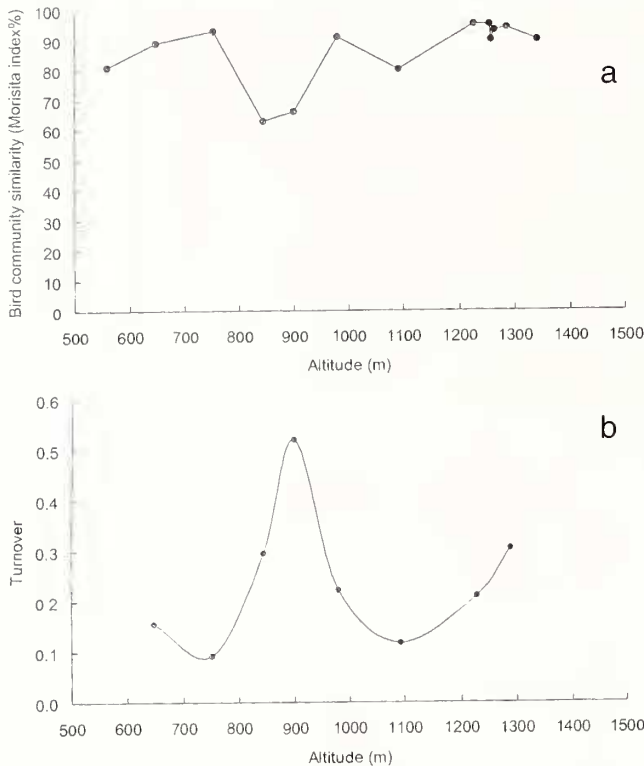


Fig. 4: Turnover in rainforest bird community composition as a function of altitude in KMTR. In the upper panel (a) similarity between each site and the immediately higher site is plotted against altitude of the lower site. The lower panel (b) uses data from 10 roughly equally-spaced sites along the gradient and plots rate of change in similarity between a site and the sites immediately lower and higher to it against altitude of the central site. The rate of change of similarity was calculated as $\frac{1}{2}[(d_1/w_1)+(d_2/w_2)]$, where d_1 and w_1 are the Morisita index dissimilarity and altitudinal distance between a site and the lower site, and d_2 and w_2 the corresponding values for the higher site.

under Rapoport's rule, hump-shaped relationships with productivity (Rosenzweig and Abramsky 1993), or mid-altitude peaks arising from geometric constraints (the mid-domain effect, Colwell and Lees 2000).

Instead, the observed pattern of bird species richness is a spatial analogue of the recent theoretical model proposed by Brown *et al.* (2001), to explain the regulation of local species richness over time in changing environments. This model suggests that species composition may vary substantially over time, but species richness, as an emergent property of ecosystems, is often regulated within narrow limits. Five conditions are necessary and sufficient for this to occur (Brown *et al.* 2001): (i) productivity, or resource availability, remains relatively constant, (ii) other abiotic or biotic factors vary, causing turnover in species composition, (iii) a regional species pool provides a source of colonists to local sites that are open systems with respect to species colonisation and

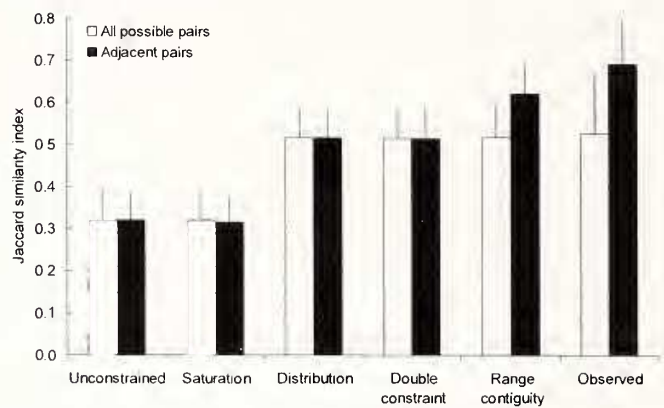


Fig. 5: Between-site similarities (average Jaccard index) in rainforest bird community composition estimated from the five non-equilibrium null models compared with observed values in the data set. Vertical lines are 1 SD.

extinction, (iv) the species pool contains species capable of utilizing the entire range of resources and showing compensatory shifts in abundance, and (v) the division rule governing apportioning of resources across species results in similar ranked species-abundance distributions.

There is partial support for these criteria for rainforest birds in KMTR. Although productivity could not be directly measured, the relative lack of clinal change in vegetation variables across the altitudinal gradient suggests that habitat structure and resource availability did not vary substantially. This is partly because a narrower altitudinal gradient was sampled in this study (500-1400 m) compared to studies from the Neotropics (Terborgh 1971, 1977; Rahbek 1997). Over a wider altitude range, bird diversity may decline with altitude and corresponding reduction in forest structural complexity as in the Peruvian Andes (Terborgh 1977). In the present study, bird species richness variables were uncorrelated to any vegetation variables except for a negative, possibly spurious, correlation with leaf litter depth. The second condition of Brown *et al.* (2001), species turnover due to changes in abiotic and biotic variables, finds clear support in the correlated variation in bird community composition in relation to altitude and tree species composition. The existence of a species pool that is nearly twice as large (with at least 58 species) as in any one local sampling site (around 30 species), adduces support for the third criterion. Finally, the occurrence of a number of species that occurred across the entire altitudinal gradient and the similar overall bird abundance and ranked species-abundance distributions (data not presented here) suggest support for the final criteria. Further support is, however, required regarding compensatory shifts in abundance of bird species along the altitudinal gradient, possibly in relation to occurrence of other bird species (Terborgh and Weske 1975).

Species range sizes and distribution

Range sizes of rainforest birds at KMTR do not conform to the expectations under Rapoport's rule, which may operate only under specific ecological conditions including competition and the rescue effect (Taylor and Gaines 1999). The failure of Rapoport's rule and the mid-domain effect is consequential to the peculiar distribution pattern of rainforest birds in KMTR. A number of species, particularly those endemic to the Western Ghats, were restricted in distribution to lower (e.g., *Ocyrceros griseus*, *Cyornis pallipes*) or higher (*Ficedula nigrorufa*, *Brachypteryx major*) altitudes (Appendix, Ali and Ripley 1983). Daniels (1992) has noted that a significant proportion of endemics, among birds and angiosperms in the Western Ghats, is restricted to the higher hills (> 1000 m). In other words, bird distributions were mainly of three types, restricted-range species of low and high altitudes, and widely distributed species, with no mid-altitude species with small ranges. The low altitude species included many that also use moist deciduous forest habitats (e.g., *Ocyrceros griseus*, *Psittacula columboides*, *Pycnonotus priocephalus*), whereas high-altitude restricted species are largely confined to wet evergreen rainforest (e.g., *Ficedula nigrorufa*, *Brachypteryx major*, *Emmyias albicaudata*, Ali and Ripley 1983). This pattern may be a consequence of historical factors that influenced the prevalence and distribution of rainforest over geological time scales.

Species turnover

A spectrum of opinion exists on the factors influencing the composition or assembly of species communities at particular sites and their variation over space and time. At one extreme, equilibrium models suggest that local communities are integrated, repeatable units whose composition is strictly regulated and predictable as a result of deterministic factors such as the varying environmental tolerances of species and competition (Clements 1916; MacArthur 1972; Pandolfi 1996; Terborgh *et al.* 1996; Pitman *et al.* 2001). In contrast, non-equilibrium approaches note that community composition varies substantially over space and time, apparently due to stochastic or historical effects of colonization and extinction (Gleason 1926; Whittaker 1970; Strong *et al.* 1984; Hubbell and Foster 1986; Brown *et al.* 2001). A fundamental distinction between non-equilibrium and equilibrium models is that under the former, community composition may be expected to 'drift' or vary continuously through space and time, whereas the latter predicts that spatially or temporally independent sites with similar environmental conditions would have similar communities (Hubbell and Foster 1986; Terborgh *et al.* 1996). Using a space-for-time substitution approach, biological influences and deterministic structure have been

demonstrated for communities of tropical rainforest trees (Terborgh *et al.* 1996; Pitman *et al.* 2001) and birds (Terborgh and Weske 1975; Raman *et al.* 1998).

The significant effect of altitude and tree species composition on bird community composition and the lack of influence of geographic distance, suggests the inapplicability of non-equilibrium models of randomly-varying distribution-abundance patterns with spatial dependence (Terborgh *et al.* 1996). Thus, high-altitude sites in Sengaltheri clustered with high-altitude sites farther away (Neterikal trail, Kakachi and Kodayar) than to virtually adjacent sites at lower altitudes. Bird community composition may be constrained by the altitude-specific environment including temperature, irradiance, and other biological factors known to vary with altitude in tropical rainforests (Richards 1996). In addition, tree species composition appears to be a key determinant of rainforest bird community composition in this study as in other studies from south-western India (Raman and Sukumar 2002), and north-eastern India (Raman *et al.* 1998).

The pattern of species turnover with altitude indicated the occurrence of distinctive community composition at low-altitudes (< 900 m) and high altitudes (> 1100 m) separated by a transitional zone of high turnover. This paralleled the observed range size distributions and placements of rainforest birds in, KMTR. The occurrence of the transition zone at around 1000-1200 m altitude may be due to environmental changes related to the formation of cloud- and mist-cover during the monsoon months in the southern Western Ghats as in other tropical montane rainforest regions (Richards 1996).

Null models of community assembly

In large-scale community studies, models with few or no constraints that attempt to parsimoniously simulate the effects of pure chance, almost invariably fail to explain community structure (Farnsworth and Ellison 1996; Terborgh *et al.* 1996; Pitman *et al.* 2001). In this study, we evaluated a hierarchy of null models with constraints that attempt to inject varying degrees of biological realism. The results clearly indicate that non-equilibrium models that do not incorporate essential biological constraints fail to predict or reproduce the observed pattern of similarities in species composition between sites. Even with constraints on site richness, species' range widths, and contiguity, the results suggest that assuming species are distributed independently of each other can result in community similarities close to, but departing significantly from, those actually observed. Thus, Whittaker's (1970) models of local community structure as a consequence of independent overlapping species distributions along ecological gradients are only partly supported. The data suggests the possibility that some species have significantly

higher overlap or avoidance due to biological determinants such as joint distributions up to ecotones or exclusive distributions due to competitive interactions (Terborgh 1971, 1985). Field research in the Western Ghats on competition and on avian distributional ecology along with null model tests (e.g. Hofer *et al.* 1999, 2000) may shed more light on these aspects.

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TROPICAL RAINFOREST BIRD COMMUNITY STRUCTURE IN WESTERN GHATS

Appendix: Range distribution and abundance (individuals/ha) of resident rainforest bird species in KMTR. Species are arranged increasing order of elevational mid-point. Rank refers to increasing correlation (Kendall Tau) between site elevation and abundance across sites A to N (see Table 1).

Scientific name	A	B	C	D	E	F	G	H	I	J	K	L	M	N	Tau	P	Rank
<i>Tephrornis gularis</i>	0.1														-0.38	0.060	17
<i>Pycnonotus priocephalus*</i>	0.1														-0.38	0.060	18
<i>Nectarinia lotenia</i>	0.1														-0.38	0.060	19
<i>Psittacula cyanocephala</i>	0.1														-0.38	0.060	20
<i>Turdoides subrufus*</i>	0.2														-0.38	0.060	21
<i>Megalaima rubricapilla</i>	0.2	0.1		0.1											-0.58	0.004	10
<i>Gracula religiosa</i>	0.5	0.3	0.1	0.6											-0.62	0.002	6
<i>Ocyrceros griseus*</i>	0.6	0.2	0.2	0.4											-0.62	0.002	7
<i>Dicurus aeneus</i>	0.1			0.1											-0.43	0.033	15
<i>Pycnonotus melanicterus</i>	0.7	0.1		0.3	0.5										-0.56	0.006	12
<i>Chloropsis aurifrons</i>	0.3	0.1	0.1		0.1										-0.64	0.001	4
<i>Dendrocitta leucogastra</i>	0.6	0.6	0.5	1.8	0.8	0.7									-0.59	0.004	9
<i>Irena puella</i>	1.0	1.1	0.5	0.9	0.6	0.8	0.8								-0.69	0.001	3
<i>Psittacula columboides*</i>	0.8	0.5	0.3	0.8	0.2	0.2	0.6								-0.71	0.000	1
<i>Hypothymis azurea</i>	0.3	0.3	0.4	0.4	0.9		0.1								-0.58	0.004	11
<i>Celeus brachyurus</i>	0.1				0.1	0.2	0.1	0.1	0.1	0.1					-0.21	0.291	28
<i>Accipiter virgatus</i>	0.2	0.1	0.1				0.1				0.1				-0.45	0.026	14
<i>Dicurus paradiseus</i>	2.0	1.1	1.1	1.2	1.1	0.4	0.5	0.4	0.2	0.7	0.5				-0.70	0.000	2
<i>Loriculus vernalis</i>	0.8	0.4		0.1		0.4	0.3		0.1		0.1	0.1			-0.43	0.034	16
<i>Cyornis pallipes*</i>	1.1	0.7	0.7	1.1	0.8	1.1	0.3		0.1		0.4	0.2	0.1		-0.53	0.009	13
<i>Chrysocolaptes lucidus</i>	0.7	0.5	0.7	0.3	0.1		0.2	0.5	0.3	0.1	0.4	0.4	0.3	0.3	-0.23	0.254	27
<i>Ducula badia</i>	0.8	1.5	1.5	1.3	0.3	0.1	0.2								-0.63	0.002	5
<i>Gallus sonneratii</i>	0.5	0.2	0.7	0.6	0.4	0.1	0.1	0.2	0.3	0.1					-0.62	0.002	8
<i>Nectarinia minima*</i>	3.1	2.8	3.2	3.3	3.0	2.7	2.9	3.2	1.2	2.8	2.9	3.1	0.4	1.2	-0.37	0.065	22
<i>Harpactes fasciatus</i>	0.3	0.5	0.3		0.2	0.2	0.3	0.3	0.2	0.1	0.2	0.1	0.1	0.2	-0.35	0.080	23
<i>Pellorneum ruficeps</i>	0.4	0.5	0.8	0.9	0.3	0.5	0.3	0.4	0.9	0.2	0.5	0.2	0.4	0.3	-0.28	0.162	24
<i>Megalaima viridis</i>	0.5	1.0	0.3	0.7	0.1	0.4	1.2				0.1	0.1	0.3	0.3	-0.27	0.171	25
<i>Iole indica</i>	1.3	2.1	2.3	1.8	2.3	2.4	2.8	1.0	0.6	0.6	1.1	1.6	1.1	1.1	-0.26	0.203	26
<i>Pericrocotus flammeus</i>	0.9	0.4	0.7	0.1	0.4	0.8	1.0	0.2	0.4		0.2	0.3	0.5	0.7	-0.12	0.543	29
<i>Sitta frontalis</i>	0.5	0.8	0.2	0.3	0.5			0.3	0.3	0.1	0.3	0.2	0.4	0.8	-0.02	0.910	33
<i>Alcippe poiocephala</i>	1.0	3.3	1.7	2.0	2.0	2.3	2.7	4.3	3.5	4.0	1.8	1.9	3.0	1.4	0.12	0.547	36
<i>Myophonus horsfieldii</i>	0.4	0.3	0.4	0.4	0.1	0.2	0.2	0.8	0.3	0.5	0.4	1.1	0.4	0.4	0.24	0.240	39
<i>Spilornis cheela</i>											0.1				0.26	0.192	40
<i>Dicaeum concolor</i>	0.1	0.3	0.5	0.4	0.3	0.3	1.0	0.6	0.4	0.3	0.1	0.6	0.7	0.9	0.33	0.104	42

Appendix (contd.): Range distribution and abundance (individuals/ha) of resident rainforest bird species in KMTR. Species are arranged increasing order of elevational mid-point. Rank refers to increasing correlation (Kendall Tau) between site elevation and abundance across sites A to N (see Table 1).

Scientific name	A	B	C	D	E	F	G	H	I	J	K	L	M	N	Tau	P	Rank
<i>Arachnothera longirostra</i>	0.1	0.2	0.2	0.2			0.3	0.9	0.4	0.7	0.2	0.5	0.4	0.5	0.46	0.023	49
<i>Pomatorhinus horsfieldii</i>	0.3	0.1	0.3	0.4	0.9	0.6	0.6	1.2	1.1	0.9	0.8	0.4	1.1	1.1	0.49	0.014	50
<i>Garullax delesserti</i> *		1.4			5.0			0.8	0.8					1.0	-0.03	0.887	32
<i>Dinopium javanense</i>			0.1	0.1		0.1		0.1	0.2	0.1	0.1				-0.01	0.944	34
<i>Dryocopus javensis</i>				0.1					0.1						-0.04	0.831	30
<i>Galloperdix spadicea</i> (a)							0.1								-0.03	0.885	31
<i>Rhopocichla atriceps</i>		0.1	0.5				0.2		0.5	0.8	0.4	0.2	0.4	0.4	0.35	0.080	43
<i>Chalcophaps indica</i>			0.1			0.1		0.2	0.2	0.1	0.1	0.1	0.1	0.1	0.36	0.072	45
<i>Lonchura kelaarti</i>			0.2											0.2	0.09	0.670	35
<i>Zoothera citrina</i>			0.1	0.1				0.2	0.2	0.4	0.4	0.1	0.1	0.2	0.55	0.006	53
<i>Hypsipetes leucocephalus</i>					0.2	0.1	0.6	0.7	1.8	1.7	0.2	0.7	1.3	1.1	0.66	0.001	56
<i>Zosterops palpebrosus</i>					0.8	1.2	3.1	6.1	5.7	5.0	6.2	11	8.9	7.6	0.83	0.000	58
<i>Hemipus picatus</i>						0.2	0.2	0.5	0.2	0.5	0.2	0.2	0.4	0.8	0.70	0.000	57
<i>Turdus merula</i> (b)							0.4	3.0	1.8	2.6	2.2	1.8	1.8	2.8	0.15	0.469	37
<i>Culicicapa ceylonensis</i>											0.1				0.64	0.001	55
<i>Ceyx erithacus</i> (c)											0.1	0.3	0.1		0.20	0.311	38
<i>Parus xanthogenys</i>								0.1	0.2	0.4	0.1	0.3	0.1		0.51	0.012	51
<i>Eumyias albicaudata</i> *								1.2	0.8	0.4	0.1	0.6	1.1	1.1	0.52	0.010	52
<i>Ficedula nigrorufa</i> *								1.8	1.2	1.9	0.8	0.2	1.4	2.5	0.63	0.002	54
<i>Garrulax jerdoni</i> *									0.4					2.2	0.36	0.076	44
<i>Picumnus innominatus</i>											0.1			0.1	0.43	0.033	48
<i>Pycnonotus jocosus</i> (d)												0.3			0.32	0.111	41
<i>Gorsachius melanolophus</i>														0.1	0.38	0.060	46
<i>Brachypteryx major</i> *														0.3	0.38	0.060	47

*Species endemic to Western Ghats. **Incomplete data:** (a) also occurs sporadically in elevations below site G, more infrequently above, (b) also occurs in higher elevation sites, (c) vagrant, seen only once during the study, (d) a non-forest species, occurring at the rainforest edge only in the slightly disturbed site M.